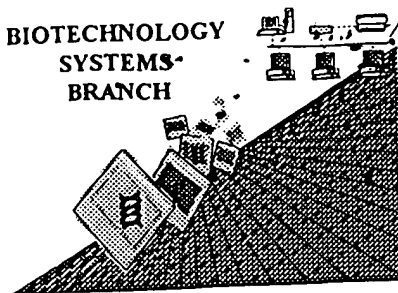


RAW SEQUENCE LISTING ERROR REPORT

BIOTECHNOLOGY
SYSTEMS
BRANCH



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:

09/829 936

Source:

OIPB

Date Processed by STIC:

10/15/01

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 - 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>

Raw Sequence Listing Error Summary

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER:

09/829936

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY P

- 1 Wrapped Nucleics
 Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2 Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.
- 3 Misaligned Amino
 Numbering The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
- 4 Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
- 5 Variable Length Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6 PatentIn 2.0
 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 7 Skipped Sequences
 (OLD RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence
 (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 This sequence is intentionally skipped

 Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
- 8 Skipped Sequences
 (NEW RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence:
 <210> sequence id number
 <400> sequence id number
 000
- 9 Use of n's or Xaa's
 (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
 Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
 In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents
- 10 Invalid <213>
 Response Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or Artificial Sequence
- 11 Use of <220> Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses.
 Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
 (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12 PatentIn 2.0
 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
- 13 Misuse of n n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.

AMC/MH - Biotechnology Systems Branch - 08/21/2001

The type of errors shown exist throughout the Sequence Listing. Please check subsequent sequences for similar errors.

Global Errors

OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/829,936

DATE: 10/15/2001

TIME: 16:16:09

Input Set : A:\ES.txt

Output Set: N:\CRF3\10152001\I829936.raw

3 <110> APPLICANT: Aventis Pharma, S.A.
 5 <120> TITLE OF INVENTION: Polyopeptide (MBP1) Capable Of Interacting With Oncogenic Mutants Of The

6 P53 Protein

8 <130> FILE REFERENCE: ST98033

10 <140> CURRENT APPLICATION NUMBER: 09/829,936

11 <141> CURRENT FILING DATE: 2001-04-11

13 <150> PRIOR APPLICATION NUMBER: FR9812754

14 <151> PRIOR FILING DATE: 1998-10-12

16 <160> NUMBER OF SEQ ID NOS: 33

18 <170> SOFTWARE: PatentIn version 3.1

20 <210> SEQ ID NO: 1

21 <211> LENGTH: 23

22 <212> TYPE: DNA

23 <213> ORGANISM: Artificial Sequence: Oligonucleotide

W--> 25 <220> FEATURE:

W--> 25 <223> OTHER INFORMATION: Errored

25 <400> SEQUENCE: 1

26 agatctgtat ggaggagccg cag

29 <210> SEQ ID NO: 2

30 <211> LENGTH: 29

31 <212> TYPE: DNA

32 <213> ORGANISM: Artificial Sequence: Oligonucleotide 3' -393 (p53)

W--> 34 <220> FEATURE:

W--> 34 <223> OTHER INFORMATION: Errored

34 <400> SEQUENCE: 2

35 agatctcatc agtctgagtc aggccttc

38 <210> SEQ ID NO: 3

39 <211> LENGTH: 15

40 <212> TYPE: DNA

41 <213> ORGANISM: Artificial Sequence: Oligonucleotide H175 3'

W--> 43 <220> FEATURE:

W--> 43 <223> OTHER INFORMATION:

43 <400> SEQUENCE: 3

44 ggggcagtgc ctcac

47 <210> SEQ ID NO: 4

48 <211> LENGTH: 15

49 <212> TYPE: DNA

50 <213> ORGANISM: Artificial Sequence: Oligonucleotide W248 3'

W--> 52 <220> FEATURE:

W--> 52 <223> OTHER INFORMATION: mandatory response

52 <400> SEQUENCE: 4

53 gggcctccag ttcac

56 <210> SEQ ID NO: 5

57 <211> LENGTH: 15

58 <212> TYPE: DNA

59 <213> ORGANISM: Artificial Sequence: Oligonucleotide H273 3'

W--> 61 <220> FEATURE:

Does Not Comply
Corrected Diskette Needed

A 213 response of "Artificial Sequence" requires an explanation of in which 223; for example, 23 oligonucleotide

29

The type of errors shown exist throughout the Sequence Listing. Please check subsequent sequences for similar errors.

15

- Global Errors

The type of errors shown exist throughout the Sequence Listing. Please check subsequent sequences for similar errors.

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/829,936

DATE: 10/15/2001

TIME: 16:16:09

Input Set : A:\ES.txt

Output Set: N:\CRF3\10152001\I829936.raw

W--> 61 <223> OTHER INFORMATION:

61 <400> SEQUENCE: 5

62 acaaacatgc acctc

15

65 <210> SEQ ID NO: 6

66 <211> LENGTH: 15

67 <212> TYPE: DNA

68 <213> ORGANISM: Artificial Sequence: Oligonucleotide G281 3'

W--> 70 <220> FEATURE:

W--> 70 <223> OTHER INFORMATION:

70 <400> SEQUENCE: 6

71 gcgccggcct ctccc

15

74 <210> SEQ ID NO: 7

75 <211> LENGTH: 23

76 <212> TYPE: DNA

77 <213> ORGANISM: Artificial Sequence: Oligonucleotide 5' -73

W--> 79 <220> FEATURE:

W--> 79 <223> OTHER INFORMATION:

79 <400> SEQUENCE: 7

80 agatctgtgt ggcacctgca cca

23

83 <210> SEQ ID NO: 8

84 <211> LENGTH: 1021

85 <212> TYPE: DNA

86 <213> ORGANISM: Artificial Sequence: Fragment C-term MBP1 murine: CDS

W--> 87 (1)..(885)

W--> 90 <220> FEATURE:

W--> 90 <223> OTHER INFORMATION:

90 <400> SEQUENCE: 8

91 tgcacctgcc ctgatggtta ccgaaaaatt ggacctgaat gtgtggacat agatgagtgt 60

93 cgttaccgct attgccagca tcgatgtgtg aacctgccgg gctcctttcg atgccagtgt 120

95 gagccaggct tccagttggg acctaaacac cgctcttctg tggatgtgaa tgagtgtgac 180

97 atgggagccc catgtgagca gcgctgcttc aactcctatg ggaccttcct gtgtcgctgt 240

99 aaccagggct atgagctgca ccgggatggc ttctcctgca gcgatatcga tgagtgcggc 300

101 tactccagtt acctctgcca gtaccgctgt gtcaacgagc caggccgatt ctctgtcac 360

103 tgcccacaag gctaccagct gctggctaca aggctctgcc aagatattga cgagtgtgaa 420

105 acaggtgcac accaatgttc tgaggcccaa acctgtgtca acttccatgg gggttaccgc 480

107 tgtgtggaca ccaaccgttg tgtggagccc tatgtccaag tgtcagacaa ccgctgcctc 540

109 tgccctgcct ccaatcccct ttgtcgagag cagccttcat ccattgtgca ccgctacatg 600

111 agcatcacct cagagcgaag tgtgcctgct gacgtgttcc agatccaggc aacctctgtc 660

113 taccctgggt cctacaatgc ctttcagatc cgttctggaa acacacaggg ggacttctac 720

115 attaggcaaa tcaacaatgt cagcgccatg ctggctcctg ccaggccagt gacgggaccc 780

117 cgggagtagc tgctggacct ggagatggtc accatgaatt cccttatgag ctaccggggc 840

119 agctctgtac tgagactcac ggtctttgtg ggagcctata ccttctgaag accctcaggg 900

121 aagggccatg tgggggcccc ttccccctcc catagcttaa gcagccccgg gggcctaggg 960

123 atgaccgttc tgcttaaagg aactatgatg tgaaggacaa taaagggaga aagaaggaaa 1020

125 a 1021

128 <210> SEQ ID NO: 9

129 <211> LENGTH: 295

130 <212> TYPE: PRT

131 <213> ORGANISM: Artificial Sequence: Fragment C-term MBP1 murine

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/829,936

DATE: 10/15/2001

TIME: 16:16:09

Input Set : A:\ES.txt

Output Set: N:\CRF3\10152001\I829936.raw

W--> 133 <220> FEATURE:

W--> 133 <223> OTHER INFORMATION:

133 <400> SEQUENCE: 9

135 Cys Thr Cys Pro Asp Gly Tyr Arg Lys Ile Gly Pro Glu Cys Val Asp

136 1 5 10 15

139 Ile Asp Glu Cys Arg Tyr Arg Tyr Cys Gln His Arg Cys Val Asn Leu

140 20 25 30

143 Pro Gly Ser Phe Arg Cys Gln Cys Glu Pro Gly Phe Gln Leu Gly Pro

144 35 40 45

147 Asn Asn Arg Ser Cys Val Asp Val Asn Glu Cys Asp Met Gly Ala Pro

148 50 55 60

151 Cys Glu Gln Arg Cys Phe Asn Ser Tyr Gly Thr Phe Leu Cys Arg Cys

152 65 70 75 80

155 Asn Gln Gly Tyr Glu Leu His Arg Asp Gly Phe Ser Cys Ser Asp Ile

156 85 90 95

159 Asp Glu Cys Gly Tyr Ser Ser Tyr Leu Cys Gln Tyr Arg Cys Val Asn

160 100 105 110

163 Glu Pro Gly Arg Phe Ser Cys His Cys Pro Gln Gly Tyr Gln Leu Leu

164 115 120 125

167 Ala Thr Arg Leu Cys Gln Asp Ile Asp Glu Cys Glu Thr Gly Ala His

168 130 135 140

171 Gln Cys Ser Glu Ala Gln Thr Cys Val Asn Phe His Gly Gly Tyr Arg

172 145 150 155 160

175 Cys Val Asp Thr Asn Arg Cys Val Glu Pro Tyr Val Gln Val Ser Asp

176 165 170 175

179 Asn Arg Cys Leu Cys Pro Ala Ser Asn Pro Leu Cys Arg Glu Gln Pro

180 180 185 190

183 Ser Ser Ile Val His Arg Tyr Met Ser Ile Thr Ser Glu Arg Ser Val

184 195 200 205

187 Pro Ala Asp Val Phe Gln Ile Gln Ala Thr Ser Val Tyr Pro Gly Ala

188 210 215 220

191 Tyr Asn Ala Phe Gln Ile Arg Ser Gly Asn Thr Gln Gly Asp Phe Tyr

192 225 230 235 240

195 Ile Arg Gln Ile Asn Asn Val Ser Ala Met Leu Val Leu Ala Arg Pro

196 245 250 255

199 Val Thr Gly Pro Arg Glu Tyr Val Leu Asp Leu Glu Met Val Thr Met

200 260 265 270

203 Asn Ser Leu Met Ser Tyr Arg Ala Ser Ser Val Leu Arg Leu Thr Val

204 275 280 285

207 Phe Val Gly Ala Tyr Thr Phe

208 290 295

211 <210> SEQ ID NO: 10

212 <211> LENGTH: 39

213 <212> TYPE: DNA

214 <213> ORGANISM: Artificial Sequence: Oligonucleotide c-myc 5'

W--> 216 <220> FEATURE:

W--> 216 <223> OTHER INFORMATION:

216 <400> SEQUENCE: 10

217 gatccatgga gcagaagctg atctccgagg aggacctga

39

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/829,936

DATE: 10/15/2001

TIME: 16:16:09

Input Set : A:\ES.txt

Output Set: N:\CRF3\10152001\I829936.raw

```

220 <210> SEQ ID NO: 11
221 <211> LENGTH: 39
222 <212> TYPE: DNA
223 <213> ORGANISM: Artificial Sequence: Oligonucleotide c-myc 3'
W--> 225 <220> FEATURE:
W--> 225 <223> OTHER INFORMATION:
225 <400> SEQUENCE: 11
226 gatctcaggt cctcctcgga gatcagcttc tgctccatg          39
229 <210> SEQ ID NO: 12
230 <211> LENGTH: 45
231 <212> TYPE: DNA
232 <213> ORGANISM: Artificial Sequence: MCS 5'
W--> 234 <220> FEATURE:
W--> 234 <223> OTHER INFORMATION:
234 <400> SEQUENCE: 12
235 gatctcggtc gacctgcatg caattcccgg gtgcggccgc gagct          45
238 <210> SEQ ID NO: 13
239 <211> LENGTH: 37
240 <212> TYPE: DNA
241 <213> ORGANISM: Artificial Sequence: MCS 3'
W--> 243 <220> FEATURE:
W--> 243 <223> OTHER INFORMATION:
243 <400> SEQUENCE: 13
244 cgcggccgca cccgggaatt gcatgcaggt cgaccga          37
247 <210> SEQ ID NO: 14
248 <211> LENGTH: 22
249 <212> TYPE: DNA
250 <213> ORGANISM: Artificial Sequence: Oligonucleotide 3' mMBP1
W--> 252 <220> FEATURE:
W--> 252 <223> OTHER INFORMATION:
252 <400> SEQUENCE: 14
253 cggtactggc agaggttaact gg
256 <210> SEQ ID NO: 15
257 <211> LENGTH: 1513
258 <212> TYPE: DNA
259 <213> ORGANISM: Artificial Sequence: MBP1 murine (complete sequence): CDS
W--> 260 (49)..(1377)
W--> 262 <220> FEATURE:
W--> 262 <223> OTHER INFORMATION:
262 <400> SEQUENCE: 15
263 gctgtggcag aaacccctga cttctgccca ccacctccca gcctcaggat gctccctttt          60
265 gcctcctgcc tccccgggtc tttgtgtctc tgggctttc tgctgttgct cttgggagca          120
267 gcgtccccac aggatcccca ggagccggac agctacacgg aatgcacaga tggctatgag          180
269 tgggatgcag acagccagca ctgccgggat gtcaacgagt gcctgaccat cccggaggct          240
271 tgcaagggtg agatgaaatg catcaaccac tacgggggtt atttgtgtct gcctcgtctc          300
273 gctgccgtca tcaagtatct ccatggtgaa ggacctccac cgccagcggc ccatgctcaa          360
275 caaccaaacc cttgcccgca gggctacgag cctgatgaac aggagagctg tgtggatgtg          420
277 gacgagtgtg cccaggcttt gcatgactgt cgccctagtc aggactgcca taaccttcct          480
279 ggctcctacc agtgcacctg ccctgatggt taccgaaaaa ttggaccgca atgtgtggac          540

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Errors

The type of errors shown exist throughout the Sequence Listing. Please check subsequent sequences for similar errors.

RAW SEQUENCE LISTING

DATE: 10/15/2001

PATENT APPLICATION: US/09/829,936

TIME: 16:16:09

Input Set : A:\ES.txt

Output Set: N:\CRF3\10152001\I829936.raw

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281 atagatgagt gtcgttaccg ctattgccag catcgatgtg tgaacctgcc gggctctttt 600
283 cgatgccagt gtgagccagg cttccagttg ggacctaac accgctcttg tgtggatgtg 660
285 aatgagtgtg acatgggagc cccatgtgag cagcgctgct tcaactccta tgggaccttc 720
287 ctgtgtcgct gtaaccaggg ctatgagctg caccgggatg gcttctcctg cagcgatctc 780
289 gatgagtgcg gctactccag ttacctctgc cagtaccgct gtgtcaacga gccaggccga 840
291 ttctcctgtc actgcccaca aggtaccag ctgctggcta caaggctctg ccaagatatt 900
293 gacgagtgtg aaacagggtg acaccaatgt tctgaggccc aaacctgtgt caacttccat 960
295 ggggggttacc gctgtgtgga caccaaccgt tgtgtggagc cctatgtcca agtgtcagac 1020
297 aaccgctgcc tctgccctgc ctccaatccc ctttgtcgag agcagccttc atccattgtg 1080
299 caccgctaca tgagcatcac ctgagagcga agtgtgcctg ctgacgtgtt tcagatccag 1140
301 gcaacctctg tctaccctgg tgctacaat gcctttcaga tccgttcttg aaacacacag 1200
303 ggggacttct acattaggca aatcaacaat gtcagcgcca tgctggctct cgccaggcca 1260
305 gtgacgggac cccgggagta cgtgctggac ctggagatgg tcaccatgaa ttcccttatg 1320
307 agctaccggg ccagctctgt actgagactc acggtctttg tgggagccta taccttctga 1380
309 agaccctcag ggaagggccca tgtggggggc ccttccccct cccatagctt aagcagcccc 1440
311 ggggggcctag ggatgaccgt tctgcttaaa ggaactatga tgtgaaggac aataaaggga 1500
313 gaaagaagga aaa 1513

```

316 <210> SEQ ID NO: 16

317 <211> LENGTH: 442

318 <212> TYPE: PRT

319 <213> ORGANISM: Artificial Sequence: MBP1 murine (complete sequence)

W--> 321 <220> FEATURE:

W--> 321 <223> OTHER INFORMATION:

321 <400> SEQUENCE: 16

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323 Met Leu Pro Phe Ala Ser Cys Leu Pro Gly Ser Leu Leu Leu Trp Ala
324 1 5 10 15
327 Phe Leu Leu Leu Leu Leu Gly Ala Ala Ser Pro Gln Asp Pro Glu Glu
328 20 25 30
331 Pro Asp Ser Tyr Thr Glu Cys Thr Asp Gly Tyr Glu Trp Asp Ala Asp
332 35 40 45
335 Ser Gln His Cys Arg Asp Tyr Asn Glu Cys Leu Thr Ile Pro Glu Ala
336 50 55 60
339 Cys Lys Gly Glu Met Lys Cys Ile Asn His Tyr Gly Gly Tyr Leu Cys
340 65 70 75 80
343 Leu Pro Arg Ser Ala Ala Val Ile Ser Asp Leu His Gly Glu Gly Pro
344 85 90 95
347 Pro Pro Pro Ala Ala His Ala Gln Gln Pro Asn Pro Cys Pro Gln Gly
348 100 105 110
351 Tyr Glu Pro Asp Glu Gln Glu Ser Cys Val Asp Val Asp Glu Cys Thr
352 115 120 125
355 Gln Ala Leu His Asp Cys Arg Pro Ser Gln Asp Cys His Asn Leu Pro
356 130 135 140
359 Gly Ser Tyr Gln Cys Thr Cys Pro Asp Gly Tyr Arg Lys Ile Gly Pro
360 145 150 155 160
363 Glu Cys Val Asp Ile Asp Glu Cys Arg Tyr Arg Tyr Cys Gln His Arg
364 165 170 175
367 Cys Val Asn Leu Pro Gly Ser Phe Arg Cys Gln Cys Glu Pro Gly Phe
368 180 185 190
371 Gln Leu Gly Pro Asn Asn Arg Ser Cys Val Asp Val Asn Glu Cys Asp

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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/829,936

DATE: 10/15/2001

TIME: 16:16:10

Input Set : A:\ES.txt

Output Set: N:\CRF3\10152001\I829936.raw

L:25 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:25 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:34 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:34 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:43 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:43 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
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L:52 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:61 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:61 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:70 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:70 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:79 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:79 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:87 M:259 W: Allowed number of lines exceeded, <213> ORGANISM:
L:90 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:90 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:133 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:133 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:216 M:258 W: Mandatory Feature missing, <220> FEATURE:
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L:225 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:225 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:234 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:234 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:243 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:243 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:252 M:258 W: Mandatory Feature missing, <220> FEATURE:
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L:260 M:259 W: Allowed number of lines exceeded, <213> ORGANISM:
L:262 M:258 W: Mandatory Feature missing, <220> FEATURE:
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L:533 M:259 W: Allowed number of lines exceeded, <213> ORGANISM:
L:535 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:535 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:590 M:258 W: Mandatory Feature missing, <220> FEATURE:
L:590 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:709 M:258 W: Mandatory Feature missing, <220> FEATURE:

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/829,936

DATE: 10/15/2001

TIME: 16:16:10

Input Set : A:\ES.txt

Output Set: N:\CRF3\10152001\I829936.raw

L:709 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:744 M:258 W: Mandatory Feature missing, <220> FEATURE:
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L:753 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:
L:797 M:259 W: Allowed number of lines exceeded, <213> ORGANISM:
L:934 M:259 W: Allowed number of lines exceeded, <213> ORGANISM: